SEQUENCE LISTING

```
<110> XIA, TAI-HE
     NI, DONGHUI
     EISHINGDRELO, HAIFENG
     ARDATI, ALI
     MINNICH, ANNE
      JUPP, RAY
<120> NOVEL G PROTEIN-COUPLED RECEPTOR
<130> 41491
<140> 09/886,041
<141> 2001-06-22
<160> 12
<170> PatentIn Ver. 2:
<210> 1
<211> 1041
<212> DNA
<213> Homo sapiens
<400> 1
atgtacaacg ggtcgtgctg ccgcatcgag ggggacacca tctcccaggt gatgccgccg 60
ctgctcattg tggcctttgt gctgggcgca ctaggcaatg gggtcgccct gtgtggtttc 120
tgcttccaca tgaagacctg gaagcccagc actgtttacc ttttcaattt ggccgtggct 180
gatttcctcc ttatgatctg cctgcctttt cggacagact attacctcag acgtagacac 240
tgggcttttg gggacattcc ctgccgagtg gggctcttca cgttggccat gaacagggcc 300
gggagcatcg tgttccttac ggtggtggct gcggacaggt atttcaaagt ggtccacccc 360
caccacgegg tgaacactat ctccaccegg gtggcggctg gcatcgtctg caccctgtgg 420
gccctggtca tcctgggaac agtgtatctt ttgctggaga accatctctg cgtgcaagag 480
acggccgtct cctgtgagag cttcatcatg gagtcggcca atggctggca tgacatcatg 540
ttccagctgg agttctttat gcccctcggc atcatcttat tttgctcctt caagattgtt 600
tggagcetga ggcggaggca gcagctggcc agacaggctc ggatgaagaa ggcgaccegg 660
ttcatcatgg tggtggcaat tgtgttcatc acatgctacc tgcccagcgt gtctgctaga 720
ctctatttcc tctggacggt gcctcgagt gcctgcgatc cctctgtcca tggggccctg 780
cacataaccc tcagcttcac ctacatgaac agcatgctgg atcccctggt gtattatttt 840
tcaagcccct cctttcccaa attctacaac aagctcaaaa tctgcagtct gaaacccaag 900
cagccaggac actcaaaaac acaaaggccg gaagagatgc caatttcgaa cctcggtcgc 960
aggagttgca tcagtgtggc aaatagtttc caaagccagt ctgatgggca atgggatccc 1020
                                                                   1041
cacattgttg agtggcactg a
<210> 2
<211> 346
<212> PRT
<213> Homo sapiens
```

Met Tyr Asn Gly Ser Cys Cys Arg Ile Glu Gly Asp Thr Ile Ser Gln

Val Met Pro Pro Leu Leu Ile Val Ala Phe Val Leu Gly Ala Leu Gly
20 25 30

<400> 2

- Asn Gly Val Ala Leu Cys Gly Phe Cys Phe His Met Lys Thr Trp Lys
 35 40 45
- Pro Ser Thr Val Tyr Leu Phe Asn Leu Ala Val Ala Asp Phe Leu Leu 50 55 60
- Met Ile Cys Leu Pro Phe Arg Thr Asp Tyr Tyr Leu Arg Arg Arg His 65 70 75 80
- Trp Ala Phe Gly Asp Ile Pro Cys Arg Val Gly Leu Phe Thr Leu Ala 85 90 95
- Met Asn Arg Ala Gly Ser Ile Val Phe Leu Thr Val Val Ala Ala Asp 100 105 110
- Arg Tyr Phe Lys Val Val His Pro His His Ala Val Asn Thr Ile Ser 115 120 125
- Thr Arg Val Ala Ala Gly Ile Val Cys Thr Leu Trp Ala Leu Val Ile 130 135 140
- Leu Gly Thr Val Tyr Leu Leu Leu Glu Asn His Leu Cys Val Gln Glu 145 150 155 160
- Thr Ala Val Ser Cys Glu Ser Phe Ile Met Glu Ser Ala Asn Gly Trp 165 170 175
- His Asp Ile Met Phe Gln Leu Glu Phe Phe Met Pro Leu Gly Ile Ile 180 185 190
- Leu Phe Cys Ser Phe Lys Ile Val Trp Ser Leu Arg Arg Arg Gln Gln
 195 200 205
- Leu Ala Arg Gln Ala Arg Met Lys Lys Ala Thr Arg Phe Ile Met Val 210 215 220
- Val Ala Ile Val Phe Ile Thr Cys Tyr Leu Pro Ser Val Ser Ala Arg 225 230 235 240
- Leu Tyr Phe Leu Trp Thr Val Pro Ser Ser Ala Cys Asp Pro Ser Val 245 250 255
- His Gly Ala Leu His Ile Thr Leu Ser Phe Thr Tyr Met Asn Ser Met 260 265 270
- Leu Asp Pro Leu Val Tyr Tyr Phe Ser Ser Pro Ser Phe Pro Lys Phe 275 280 285
- Tyr Asn Lys Leu Lys Ile Cys Ser Leu Lys Pro Lys Gln Pro Gly His 290 295 300
- Ser Lys Thr Gln Arg Pro Glu Glu Met Pro Ile Ser Asn Leu Gly Arg 305 310 315 320
- Arg Ser Cys Ile Ser Val Ala Asn Ser Phe Gln Ser Gln Ser Asp Gly
 325 330 335

Gln Trp Asp Pro His Ile Val Glu Trp His 340 345

<213> Artificial Sequence

<210> 3 <211> 20 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Primer <400> 3 ctatttcctc tggacggtgc 20 <210> 4 <211> 19 <212> DNA <213> Artificial Sequence <223> Description of Artificial Sequence: Primer <400> 4 ttatgtgcag ggccccatg 19 <210> 5 <211> 39 <212> DNA <213> Artificial Sequence <223> Description of Artificial Sequence: Primer <400> 5 tcatgtggaa gcagaaacca cacagggcga ccccattgc 39 <210> 6 <211> 21 <212> DNA <213> Artificial Sequence <223> Description of Artificial Sequence: Primer <400> 6 atgtacaacg ggtcgtgctg c 21 <210> 7 <211> 22 <212> DNA

```
<220>
<223> Description of Artificial Sequence: Primer
<400> 7
                                                                    22
tcagtgccac tcaacaatgt gg
<210> 8
<211> 20
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 8
                                                                    20
taatacgact cactataggg
<210> 9
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
                                                                    20
cagtaaacag ctatgaccat
<210> 10
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 10
ctatttcctc tggacggtgc
                                                                    20
<210> 11
<211> 19
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 11
ttatgtgcag ggccccatg
                                                                    19
<210> 12
<211> 23
```

<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Probe
<400> 12
tcgagtgcct gcgatccctc tgt